

#16
DMT
5-16-01

BATCH

RAW SEQUENCE LISTING
PATENT APPLICATION: US/09/665,728

DATE: 03/29/2001
TIME: 10:39:11

Input Set : A:\SCIOS.txt
Output Set: N:\CRF3\03292001\I665728.raw

ENTERED

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4 <110> APPLICANT: Stanton, Lawrence W.
5 Kapoun, Ann Marie
7 <120> TITLE OF INVENTION: SECRETED FACTORS
11 <130> FILE REFERENCE: SCIOS.013A
C--> 13 <140> CURRENT APPLICATION NUMBER: US/09/665,728
C--> 13 <141> CURRENT FILING DATE: 2000-09-20
13 <150> PRIOR APPLICATION NUMBER: 60/156,277
14 <151> PRIOR FILING DATE: 1999-09-27
16 <160> NUMBER OF SEQ ID NOS: 19
18 <170> SOFTWARE: FastSEQ for Windows Version 4.0
20 <210> SEQ ID NO: 1
21 <211> LENGTH: 275
22 <212> TYPE: PRT
23 <213> ORGANISM: Rattus norvegicus
25 <400> SEQUENCE: 1
26 Met Thr Pro Arg Ala Gln Leu Leu Pro Leu Leu Ala Thr Tyr Thr
27 1 5 10 15
28 Val Val Ala Ala Ala Val Thr Ser Asp Glu Pro Thr Lys Thr Leu Ser
29 20 25 30
30 Pro Ala Thr Gly Asp Ala Thr Leu Ala Phe Val Phe Asp Val Thr Gly
31 35 40 45
32 Ser Met Trp Asp Asp Leu Met Gln Val Ile Asp Gly Ala Ser Arg Ile
33 50 55 60
34 Leu Glu Arg Ser Leu Ser Ser Arg Ser Arg Val Ile Ala Asn Tyr Ala
35 65 70 75 80
36 Leu Val Pro Phe His Asp Pro Asp Ile Gly Pro Val Thr Leu Thr Ala
37 85 90 95
38 Asp Pro Val Val Phe Gln Arg Glu Leu Arg Gln Leu Tyr Val Gln Gly
39 100 105 110
40 Gly Gly Asp Cys Pro Glu Met Ser Val Gly Ala Ile Lys Ala Ala Val
41 115 120 125
42 Glu Val Ala Asn Pro Gly Ser Phe Ile Tyr Val Phe Ser Asp Ala Arg
43 130 135 140
44 Ala Lys Asp Tyr His Lys Asn Glu Leu Leu Gln Leu Leu Gln Leu
45 145 150 155 160
46 Lys Gln Ser Gln Val Val Phe Val Leu Thr Gly Asp Cys Gly Asp Arg
47 165 170 175
48 Thr His Pro Gly Tyr Leu Ala Phe Glu Glu Ile Ala Ser Thr Ser Ser
49 180 185 190
50 Gly Gln Val Phe Gln Leu Asp Lys Gln Gln Val Ser Glu Val Leu Lys
51 195 200 205
52 Trp Val Glu Ser Ala Ile Gln Ala Ser Lys Val His Leu Leu Ser Ala
53 210 215 220
54 Asp His Glu Glu Glu Gly Glu His Thr Trp Arg Ile Pro Phe Asp Pro
55 225 230 235 240
56 Ser Leu Lys Glu Val Thr Ile Ser Leu Ser Gly Pro Gly Pro Glu Ile
57 245 250 255

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58 Glu Val Arg Asp Pro Leu Gly Met Ser Gln Gly Ser Pro Pro Leu Leu
59      260          265          270
60 Met Gln Asp
61      275
64 <210> SEQ ID NO: 2
65 <211> LENGTH: 1031
66 <212> TYPE: DNA
67 <213> ORGANISM: Rattus norvegicus
69 <220> FEATURE:
70 <221> NAME/KEY: CDS
71 <222> LOCATION: (96)...(920)
73 <400> SEQUENCE: 2
74 tctagcgaac cccttcggcc cgctagagcg agactgcact gccatctatc cctgcgacct 60
75 ggcgtccca ttagggctgc agcctccggc tcagc atg acg cct agg gcg cag    113
76                               Met Thr Pro Arg Ala Gln
77                               1           5
79 ctc ctg ccg ctg ctc ctg gcg acc tac aca gta gtg gcg gcg gtc   161
80 Leu Leu Pro Leu Leu Ala Thr Tyr Thr Val Val Ala Ala Ala Val
81      10          15          20
83 aca tct gat gag ccc acg aag acg ctg tcc ccc gcc aca gga gac gcc  209
84 Thr Ser Asp Glu Pro Thr Lys Thr Leu Ser Pro Ala Thr Gly Asp Ala
85      25          30          35
87 acc ctg gcc ttc gtc ttc gat gtc acc ggc tcc atg tgg gac gat ctg  257
88 Thr Leu Ala Phe Val Phe Asp Val Thr Gly Ser Met Trp Asp Asp Leu
89      40          45          50
91 atg cag gtg atc gac ggc gcc tca cgc att ctg gag cgc agt ctg agc  305
92 Met Gln Val Ile Asp Gly Ala Ser Arg Ile Leu Glu Arg Ser Leu Ser
93      55          60          65          70
95 agc cgc agc cgg gtc atc gcc aac tat gcg ctg gtg cct ttc cac gac  353
96 Ser Arg Ser Arg Val Ile Ala Asn Tyr Ala Leu Val Pro Phe His Asp
97      75          80          85
99 cca gac att ggc cca gtg acc ctc acg gcg gac cca gtg gtg ttt cag  401
100 Pro Asp Ile Gly Pro Val Thr Leu Thr Ala Asp Pro Val Val Phe Gln
101     90          95          100
103 aga gag ctg aga caa ctc tat gtt cag gga ggt ggt gac tgc cca gaa  449
104 Arg Glu Leu Arg Gln Leu Tyr Val Gln Gly Gly Asp Cys Pro Glu
105     105         110         115
107 atg agt gtg ggg gcc atc aag gct gcc gtg gag gtt gcc aac ccc ggc  497
108 Met Ser Val Gly Ala Ile Lys Ala Ala Val Glu Val Ala Asn Pro Gly
109     120         125         130
111 tcc ttc atc tac gtc ttc tcg gat gcc cgt gcc aag gac tac cac aag  545
112 Ser Phe Ile Tyr Val Phe Ser Asp Ala Arg Ala Lys Asp Tyr His Lys
113     135         140         145         150
115 aag aat gag ctc ctg cag ctc ctg cag tcg aag cag tcg cag gtg gtc  593
116 Lys Asn Glu Leu Leu Gln Leu Leu Gln Leu Lys Gln Ser Gln Val Val
117     155         160         165
119 ttc gtg ctg act ggg gac tgc ggt gac cgc acc cac cct ggc tac ctg  641
120 Phe Val Leu Thr Gly Asp Cys Gly Asp Arg Thr His Pro Gly Tyr Leu
121     170         175         180

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123 gct ttt gag gag atc gcc tcc acc agt tct ggc caa gtg ttc cag ctg 689
124 Ala Phe Glu Glu Ile Ala Ser Thr Ser Ser Gly Gln Val Phe Gln Leu
125 185 190 195
127 gac aag cag cag gtg tcg gag gtg tta aag tgg gtg gag tcc gcc atc 737
128 Asp Lys Gln Gln Val Ser Glu Val Leu Lys Trp Val Glu Ser Ala Ile
129 200 205 210
131 cag gcc tcc aaa gtt cat ctg ctg tca gca gac cac gag gag ggc 785
132 Gln Ala Ser Lys Val His Leu Leu Ser Ala Asp His Glu Glu Glu Gly
133 215 220 225 230
135 gaa cac aca tgg aga atc cct ttt gac ccc agc ttg aag gaa gtc acc 833
136 Glu His Thr Trp Arg Ile Pro Phe Asp Pro Ser Leu Lys Glu Val Thr
137 235 240 245
139 atc tca ctg agc ggg cca ggg cct gag atc gaa gtc cgg gac cca ctg 881
140 Ile Ser Leu Ser Gly Pro Gly Pro Glu Ile Glu Val Arg Asp Pro Leu
141 250 255 260
143 ggt atg tcc cag ggt tca cct ctt ctg atg caa gac tgagctggaa 930
144 Gly Met Ser Gln Gly Ser Pro Pro Leu Leu Met Gln Asp
145 265 270 275
147 ggccaggctg aggcgatgga aggaggggcc tgaggagatg gctcagccaa taaaatgtct 990
148 gcctcacaca aaaaaaaaaa aagccggct cgagcggccg c 1031
150 <210> SEQ ID NO: 3
151 <211> LENGTH: 25
152 <212> TYPE: DNA
153 <213> ORGANISM: Artificial Sequence
155 <220> FEATURE:
156 <223> OTHER INFORMATION: synthetic
158 <400> SEQUENCE: 3
159 cgtatgttgt gtggaatgt gagcg 25
161 <210> SEQ ID NO: 4
162 <211> LENGTH: 25
163 <212> TYPE: DNA
164 <213> ORGANISM: Artificial Sequence
166 <220> FEATURE:
167 <223> OTHER INFORMATION: synthetic
169 <400> SEQUENCE: 4
170 gatgtgctgc aaggcgattt agttt 25
172 <210> SEQ ID NO: 5
173 <211> LENGTH: 28
174 <212> TYPE: DNA
175 <213> ORGANISM: Artificial Sequence
177 <220> FEATURE:
178 <223> OTHER INFORMATION: synthetic
180 <400> SEQUENCE: 5
181 gccggccagtg tgctggattt cggcttgc 28
183 <210> SEQ ID NO: 6
184 <211> LENGTH: 28
185 <212> TYPE: DNA
186 <213> ORGANISM: Artificial Sequence
188 <220> FEATURE:

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Input Set : A:\SCIOS.txt
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189 <223> OTHER INFORMATION: synthetic
191 <400> SEQUENCE: 6
192 cgaattctgc agatattccat cacactgg 28
194 <210> SEQ ID NO: 7
195 <211> LENGTH: 25
196 <212> TYPE: DNA
197 <213> ORGANISM: Artificial Sequence
199 <220> FEATURE:
200 <223> OTHER INFORMATION: synthetic
202 <400> SEQUENCE: 7
203 ctagaggccc caattcgccc tata 25
205 <210> SEQ ID NO: 8
206 <211> LENGTH: 25
207 <212> TYPE: DNA
208 <213> ORGANISM: Artificial Sequence
210 <220> FEATURE:
211 <223> OTHER INFORMATION: synthetic
213 <400> SEQUENCE: 8
214 tgagtcttat tacaattcac tggcc 25
216 <210> SEQ ID NO: 9
217 <211> LENGTH: 20
218 <212> TYPE: DNA
219 <213> ORGANISM: Artificial Sequence
221 <220> FEATURE:
222 <223> OTHER INFORMATION: synthetic
224 <400> SEQUENCE: 9
225 gctcggatcc actagtaacg 20
227 <210> SEQ ID NO: 10
228 <211> LENGTH: 18
229 <212> TYPE: DNA
230 <213> ORGANISM: Artificial Sequence
232 <220> FEATURE:
233 <223> OTHER INFORMATION: synthetic
235 <400> SEQUENCE: 10
236 tttttttttt tttttttt 18
238 <210> SEQ ID NO: 11
239 <211> LENGTH: 25
240 <212> TYPE: DNA
241 <213> ORGANISM: Artificial Sequence
243 <220> FEATURE:
244 <223> OTHER INFORMATION: synthetic
246 <400> SEQUENCE: 11
247 cgtatgttgt gtggaaattgt gagcg 25
249 <210> SEQ ID NO: 12
250 <211> LENGTH: 25
251 <212> TYPE: DNA
252 <213> ORGANISM: Artificial Sequence
254 <220> FEATURE:
255 <223> OTHER INFORMATION: synthetic

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Input Set : A:\SCIOS.txt
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257 <400> SEQUENCE: 12
258 gatgtgctgc aaggcgatta agttg 25
260 <210> SEQ ID NO: 13
261 <211> LENGTH: 20
262 <212> TYPE: DNA
263 <213> ORGANISM: Artificial Sequence
265 <220> FEATURE:
266 <223> OTHER INFORMATION: synthetic
268 <400> SEQUENCE: 13
269 tggccttcgt cttcgatgtc 20
271 <210> SEQ ID NO: 14
272 <211> LENGTH: 18
273 <212> TYPE: DNA
274 <213> ORGANISM: Artificial Sequence
276 <220> FEATURE:
277 <223> OTHER INFORMATION: synthetic
279 <400> SEQUENCE: 14
280 gccgtcgatc acctgcat 18
282 <210> SEQ ID NO: 15
283 <211> LENGTH: 22
284 <212> TYPE: DNA
285 <213> ORGANISM: Artificial Sequence
287 <220> FEATURE:
288 <223> OTHER INFORMATION: synthetic
290 <400> SEQUENCE: 15
291 ccggctccat gtgggacgat ct 22
293 <210> SEQ ID NO: 16
294 <211> LENGTH: 20
295 <212> TYPE: DNA
296 <213> ORGANISM: Artificial Sequence
298 <220> FEATURE:
299 <223> OTHER INFORMATION: synthetic
301 <400> SEQUENCE: 16
302 cggttaccac atccaagaa 20
304 <210> SEQ ID NO: 17
305 <211> LENGTH: 18
306 <212> TYPE: DNA
307 <213> ORGANISM: Artificial Sequence
309 <220> FEATURE:
310 <223> OTHER INFORMATION: synthetic
312 <400> SEQUENCE: 17
313 gcttggattt ccggggct 18
315 <210> SEQ ID NO: 18
316 <211> LENGTH: 22
317 <212> TYPE: DNA
318 <213> ORGANISM: Artificial Sequence
320 <220> FEATURE:
321 <223> OTHER INFORMATION: synthetic
323 <400> SEQUENCE: 18

VERIFICATION SUMMARY DATE: 03/29/2001
PATENT APPLICATION: US/09/665,728 TIME: 10:39:12

Input Set : A:\SCIOS.txt
Output Set: N:\CRF3\03292001\I665728.raw

L:13 M:270 C: Current Application Number differs, Replaced Current Application No
L:13 M:271 C: Current Filing Date differs, Replaced Current Filing Date